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TECH CENTER 1600/29

1600

RAW SEQUENCE LISTING   DATE: 07/09/2003
PATENT APPLICATION: US/10/042,417A   TIME: 09:07:02

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Output Set: N:\CRF4\07092003\J042417A.raw

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     3 <110> APPLICANT: Pagano, M.
     5 <120> TITLE OF INVENTION: METHODS TO IDENTIFY COMPOUNDS USEFUL FOR THE TREATMENT OF
PROLIFERATIVE
     6        AND DIFFERENTIATIVE DISORDERS
     8 <130> FILE REFERENCE: 5914-090-999
C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/042,417A
    11 <141> CURRENT FILING DATE: 2002-01-07
    13 <150> PRIOR APPLICATION NUMBER: 60/260,179
    14 <151> PRIOR FILING DATE: 2001-01-05
    16 <160> NUMBER OF SEQ ID NOS: 92
    18 <170> SOFTWARE: PatentIn Ver. 2.0
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    23 <213> ORGANISM: Homo sapiens
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P 6

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87 Glu Lys Glu Leu Cys Val Lys Tyr Phe Glu Gln Trp Ser Glu Ser Asp
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103 Lys Glu Trp Tyr Arg Val Thr Ser Asp Gly Met Leu Trp Lys Lys Leu
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106 Ile Glu Arg Met Val Arg Thr Asp Ser Leu Trp Arg Gly Leu Ala Glu
107         195                 200                 205
109 Arg Arg Gly Trp Gly Gln Tyr Leu Phe Lys Asn Lys Pro Pro Asp Gly
110     210                 215                 220
112 Asn Ala Pro Pro Asn Ser Phe Tyr Arg Ala Leu Tyr Pro Lys Ile Ile
113 225                 230                 235                 240
115 Gln Asp Ile Glu Thr Ile Glu Ser Asn Trp Arg Cys Gly Arg His Ser
116                 245                 250                 255
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119             260                 265                 270
121 Leu Gln Tyr Asp Asp Gln Lys Ile Val Ser Gly Leu Arg Asp Asn Thr
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Output Set: N:\CRF4\07092003\J042417A.raw

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133 Gly Glu Met Leu Asn Thr Leu Ile His His Cys Glu Ala Val Leu His
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136 Leu Arg Phe Asn Asn Gly Met Met Val Thr Cys Ser Lys Asp Arg Ser
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139 Ile Ala Val Trp Asp Met Ala Ser Pro Thr Asp Ile Thr Leu Arg Arg
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142 Val Leu Val Gly His Arg Ala Ala Val Asn Val Val Asp Phe Asp Asp
143 385                 390                 395                 400
145 Lys Tyr Ile Val Ser Ala Ser Gly Asp Arg Thr Ile Lys Val Trp Asn
146                 405                 410                 415
148 Thr Ser Thr Cys Glu Phe Val Arg Thr Leu Asn Gly His Lys Arg Gly
149             420                 425                 430
151 Ile Ala Cys Leu Gln Tyr Arg Asp Arg Leu Val Val Ser Gly Ser Ser
152         435                 440                 445
154 Asp Asn Thr Ile Arg Leu Trp Asp Ile Glu Cys Gly Ala Cys Leu Arg
155     450                 455                 460
157 Val Leu Glu Gly His Glu Glu Leu Val Arg Cys Ile Arg Phe Asp Asn
158 465                 470                 475                 480
160 Lys Arg Ile Val Ser Gly Ala Tyr Asp Gly Lys Ile Lys Val Trp Asp
161                 485                 490                 495
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166 Arg Thr Leu Val Glu His Ser Gly Arg Val Phe Arg Leu Gln Phe Asp
167         515                 520                 525
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191 aaggtttatt tgaaggctat tttgagaatg aagcaactgg aggaccatga agcctttgaa 420
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196 cactttcggg ggcacacggg ggcggtattt agcgtggact acaatgatga actggatatc 720
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230  65                  70                  75                  80
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233              85                  90                  95
235 Val Trp Gln Thr Ala Cys Lys Asn Leu Gly Trp Gln Ile Asp Asp Ser
236             100                 105                 110
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239         115                 120                 125
241 Arg Met Lys Gln Leu Glu Asp His Glu Ala Phe Glu Thr Ser Ser Leu
242     130                 135                 140
244 Ile Gly His Ser Ala Arg Val Tyr Ala Leu Tyr Tyr Lys Asp Gly Leu
245 145                 150                 155                 160
247 Leu Cys Thr Gly Ser Asp Asp Leu Ser Ala Lys Leu Trp Asp Val Ser
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250 Thr Gly Gln Cys Val Tyr Gly Ile Gln Thr His Thr Cys Ala Ala Val
251             180                 185                 190
253 Lys Phe Asp Glu Gln Lys Leu Val Thr Gly Ser Phe Asp Asn Thr Val
254         195                 200                 205
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260 225                   230                   235                   240
262 Leu Val Ser Gly Ser Ala Asp Phe Thr Val Lys Val Trp Ala Leu Ser
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265 Ala Gly Thr Cys Leu Asn Thr Leu Thr Gly His Thr Glu Trp Val Thr
266                   260                   265                   270
268 Lys Val Val Leu Gln Lys Cys Lys Val Lys Ser Leu Leu His Ser Pro
269             275                   280                   285
271 Gly Asp Tyr Ile Leu Leu Ser Ala Asp Lys Tyr Glu Ile Lys Ile Trp
272    290                   295                   300
275 Pro Ile Gly Arg Glu Ile Asn Cys Lys Cys Leu Lys Thr Leu Ser Val
276 305                   310                   315                   320
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279                   325                   330                   335
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287 Leu Ala Leu Leu Gly Phe Gly Asp Ile Phe Ala Leu Leu Phe Asp Asn
288    370                   375                   380
290 Arg Tyr Leu Tyr Ile Met Asp Leu Arg Thr Glu Ser Leu Ile Ser Arg
291 385                   390                   395                   400
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294                   405                   410                   415
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Output Set: **N:\CRF4\07092003\J042417A.raw**

<u>Please Note:</u>
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

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Seq#:30; Xaa Pos. 15,22,28,89
Seq#:37; N Pos. 45,329,332
Seq#:38; Xaa Pos. 110,111
Seq#:51; N Pos. 1733
Seq#:52; Xaa Pos. 576,586
Seq#:53; N Pos. 348
Seq#:54; Xaa Pos. 150,309,340,374
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